



IFWO

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/809,689

DATE: 09/03/2004  
TIME: 11:39:25

Input Set : A:\5538-1010seqlisting.txt  
Output Set: N:\CRF4\09032004\J809689.raw

3 <110> APPLICANT: Eric Potter Clarkson  
5 <120> TITLE OF INVENTION: Methods and compositions for desensitisation  
7 <130> FILE REFERENCE: 5538/1010  
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/809,689  
C--> 10 <141> CURRENT FILING DATE: 2004-03-25  
12 <150> PRIOR APPLICATION NUMBER: PCT/GB99/00080  
13 <151> PRIOR FILING DATE: 1999-01-11  
15 <150> PRIOR APPLICATION NUMBER: GB/9800445.0  
16 <151> PRIOR FILING DATE: 1998-01-09  
18 <150> PRIOR APPLICATION NUMBER: GB/9820474.6  
19 <151> PRIOR FILING DATE: 1998-09-21  
21 <160> NUMBER OF SEQ ID NOS: 124  
23 <170> SOFTWARE: PatentIn version 3.0  
25 <210> SEQ ID NO: 1  
26 <211> LENGTH: 17  
27 <212> TYPE: PRT  
28 <213> ORGANISM: Felis catus  
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32 Leu Phe Leu Thr Gly Thr Pro Asp Glu Tyr Val Glu Gln Val Ala Gln  
33 1 5 10 15  
35 Tyr  
38 <210> SEQ ID NO: 2  
39 <211> LENGTH: 16  
40 <212> TYPE: PRT  
41 <213> ORGANISM: Felis catus  
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45 Glu Gln Val Ala Gln Tyr Lys Ala Leu Pro Val Val Leu Glu Asn Ala  
46 1 5 10 15  
48 <210> SEQ ID NO: 3  
49 <211> LENGTH: 17  
50 <212> TYPE: PRT  
51 <213> ORGANISM: Felis catus  
53 <400> SEQUENCE: 3  
55 Lys Ala Leu Pro Val Val Leu Glu Asn Ala Arg Ile Leu Lys Asn Cys  
56 1 5 10 15  
58 Val  
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62 <211> LENGTH: 70  
63 <212> TYPE: PRT  
64 <213> ORGANISM: Felis catus  
66 <400> SEQUENCE: 4  
68 Glu Ile Cys Pro Ala Val Lys Asp Arg Val Asp Leu Phe Leu Thr Gly  
69 1 5 10 15

ENTERED

(pg. 6)

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71 Thr Pro Asp Glu Tyr Val Glu Gln Val Ala Gln Tyr Lys Ala Leu Pro  
72 20 25 30  
74 Val Val Leu Glu Asn Ala Arg Ile Leu Lys Asn Cys Val Asp Ala Lys  
75 35 40 45  
77 Met Thr Glu Glu Asp Lys Glu Asn Ala Leu Ser Leu Leu Asp Lys Ile  
78 50 55 60  
80 Tyr Thr Ser Pro Leu Cys  
81 65 70  
83 <210> SEQ ID NO: 5  
84 <211> LENGTH: 92  
85 <212> TYPE: PRT  
86 <213> ORGANISM: Felis catus  
88 <400> SEQUENCE: 5  
90 Val Lys Met Ala Glu Thr Cys Pro Ile Phe Tyr Asp Val Phe Phe Ala  
91 1 5 10 15  
93 Val Ala Asn Gly Asn Glu Leu Leu Leu Lys Leu Ser Leu Thr Lys Val  
94 20 25 30  
96 Asn Ala Thr Glu Pro Glu Arg Thr Ala Met Lys Lys Ile Gln Asp Cys  
97 35 40 45  
99 Tyr Val Glu Asn Gly Leu Ile Ser Arg Val Leu Asp Gly Leu Val Met  
100 50 55 60  
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110 <212> TYPE: PRT  
111 <213> ORGANISM: Felis catus  
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116 1 5 10 15  
118 Thr  
121 <210> SEQ ID NO: 7  
122 <211> LENGTH: 17  
123 <212> TYPE: PRT  
124 <213> ORGANISM: Felis catus  
126 <400> SEQUENCE: 7  
128 Arg Ile Leu Lys Asn Cys Val Asp Ala Lys Met Thr Glu Glu Asp Lys  
129 1 5 10 15  
131 Glu  
134 <210> SEQ ID NO: 8  
135 <211> LENGTH: 16  
136 <212> TYPE: PRT  
137 <213> ORGANISM: Felis catus  
139 <400> SEQUENCE: 8  
141 Lys Met Thr Glu Glu Asp Lys Glu Asn Ala Leu Ser Leu Leu Asp Lys  
142 1 5 10 15  
144 <210> SEQ ID NO: 9

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Input Set : A:\5538-1010seqlisting.txt  
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145 <211> LENGTH: 16  
146 <212> TYPE: PRT  
147 <213> ORGANISM: Felis catus  
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156 <212> TYPE: PRT  
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162 1 5 10 15  
164 <210> SEQ ID NO: 11  
165 <211> LENGTH: 17  
166 <212> TYPE: PRT  
167 <213> ORGANISM: Felis catus  
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172 1 5 10 15  
174 Leu  
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178 <211> LENGTH: 16  
179 <212> TYPE: PRT  
180 <213> ORGANISM: Felis catus  
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185 1 5 10 15  
187 <210> SEQ ID NO: 13  
188 <211> LENGTH: 16  
189 <212> TYPE: PRT  
190 <213> ORGANISM: Felis catus  
192 <400> SEQUENCE: 13  
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195 1 5 10 15  
197 <210> SEQ ID NO: 14  
198 <211> LENGTH: 16  
199 <212> TYPE: PRT  
200 <213> ORGANISM: Felis catus  
202 <400> SEQUENCE: 14  
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205 1 5 10 15  
207 <210> SEQ ID NO: 15  
208 <211> LENGTH: 16  
209 <212> TYPE: PRT  
210 <213> ORGANISM: Felis catus  
212 <400> SEQUENCE: 15  
214 Cys Tyr Val Glu Asn Gly Leu Ile Ser Arg Val Leu Asp Gly Leu Val  
215 1 5 10 15

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Input Set : A:\5538-1010seqlisting.txt  
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217 <210> SEQ ID NO: 16  
 218 <211> LENGTH: 16  
 219 <212> TYPE: PRT  
 220 <213> ORGANISM: Felis catus  
 222 <400> SEQUENCE: 16  
 224 Ser Arg Val Leu Asp Gly Leu Val Met Thr Thr Ile Ser Ser Ser Lys  
 225 1 5 10 15  
 227 <210> SEQ ID NO: 17  
 228 <211> LENGTH: 16  
 229 <212> TYPE: PRT  
 230 <213> ORGANISM: Felis catus  
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 235 1 5 10 15  
 237 <210> SEQ ID NO: 18  
 238 <211> LENGTH: 16  
 239 <212> TYPE: PRT  
 240 <213> ORGANISM: Felis catus  
 242 <400> SEQUENCE: 18  
 244 Ala Val Gln Asn Thr Val Glu Asp Leu Lys Leu Asn Thr Leu Gly Arg  
 245 1 5 10 15  
 247 <210> SEQ ID NO: 19  
 248 <211> LENGTH: 320  
 249 <212> TYPE: PRT  
 250 <213> ORGANISM: Dermatophagooides pteronyssinus  
 252 <400> SEQUENCE: 19  
 254 Met Lys Ile Val Leu Ala Ile Ala Ser Leu Leu Ala Leu Ser Ala Val  
 255 1 5 10 15  
 257 Tyr Ala Arg Pro Ser Ser Ile Lys Thr Phe Glu Glu Tyr Lys Lys Ala  
 258 20 25 30  
 260 Phe Asn Lys Ser Tyr Ala Thr Phe Glu Asp Glu Glu Ala Ala Arg Lys  
 261 35 40 45  
 263 Asn Phe Leu Glu Ser Val Lys Tyr Val Gln Ser Asn Gly Gly Ala Ile  
 264 50 55 60  
 266 Asn His Leu Ser Asp Leu Ser Leu Asp Glu Phe Lys Asn Arg Phe Leu  
 267 65 70 75 80  
 269 Met Ser Ala Glu Ala Phe Glu His Leu Lys Thr Gln Phe Asp Leu Asn  
 270 85 90 95  
 272 Ala Glu Thr Asn Ala Cys Ser Ile Asn Gly Asn Ala Pro Ala Glu Ile  
 273 100 105 110  
 275 Asp Leu Arg Gln Met Arg Thr Val Thr Pro Ile Arg Met Gln Gly Gly  
 276 115 120 125  
 278 Cys Gly Ser Cys Trp Ala Phe Ser Gly Val Ala Ala Thr Glu Ser Ala  
 279 130 135 140  
 281 Tyr Leu Ala Tyr Arg Asn Gln Ser Leu Asp Leu Ala Glu Gln Glu Leu  
 282 145 150 155 160  
 284 Val Asp Cys Ala Ser Gln His Gly Cys His Gly Asp Thr Ile Pro Arg  
 285 165 170 175  
 287 Gly Ile Glu Tyr Ile Gln His Asn Gly Val Val Gln Glu Ser Tyr Tyr

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288 180 185 190  
 290 Arg Tyr Val Ala Arg Glu Gln Ser Cys Arg Arg Pro Asn Ala Gln Arg  
 291 195 200 205  
 293 Phe Gly Ile Ser Asn Tyr Cys Gln Ile Tyr Pro Pro Asn Val Asn Lys  
 294 210 215 220  
 296 Ile Arg Glu Ala Leu Ala Gln Thr His Ser Ala Ile Ala Val Ile Ile  
 297 225 230 235 240  
 299 Gly Ile Lys Asp Leu Asp Ala Phe Arg His Tyr Asp Gly Arg Thr Ile  
 300 245 250 255  
 302 Ile Gln Arg Asp Asn Gly Tyr Gln Pro Asn Tyr His Ala Val Asn Ile  
 303 260 265 270  
 305 Val Gly Tyr Ser Asn Ala Gln Gly Val Asp Tyr Trp Ile Val Arg Asn  
 306 275 280 285  
 308 Ser Trp Asp Thr Asn Trp Gly Asp Asn Gly Tyr Gly Tyr Phe Ala Ala  
 309 290 295 300  
 311 Asn Ile Asp Leu Met Met Ile Glu Glu Tyr Pro Tyr Val Val Ile Leu  
 312 305 310 315 320  
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 315 <211> LENGTH: 146  
 316 <212> TYPE: PRT  
 317 <213> ORGANISM: Dermatophagoides pteronyssinus  
 319 <400> SEQUENCE: 20  
 321 Met Met Tyr Lys Ile Leu Cys Leu Ser Leu Leu Val Ala Ala Val Ala  
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 324 Arg Asp Gln Val Asp Val Lys Asp Cys Ala Asn His Glu Ile Lys Lys  
 325 20 25 30  
 327 Val Leu Val Pro Gly Cys His Gly Ser Glu Pro Cys Ile Ile His Arg  
 328 35 40 45  
 330 Gly Lys Pro Phe Gln Leu Glu Ala Val Phe Glu Ala Asn Gln Asn Thr  
 331 50 55 60  
 333 Lys Thr Ala Lys Ile Glu Ile Lys Ala Ser Ile Asp Gly Leu Glu Val  
 334 65 70 75 80  
 336 Asp Val Pro Gly Ile Asp Pro Asn Ala Cys His Tyr Met Lys Cys Pro  
 337 85 90 95  
 339 Leu Val Lys Gly Gln Gln Tyr Asp Ile Lys Tyr Thr Trp Asn Val Pro  
 340 100 105 110  
 342 Lys Ile Ala Pro Lys Ser Glu Asn Val Val Val Thr Val Lys Val Met  
 343 115 120 125  
 345 Gly Asp Asp Gly Val Leu Ala Cys Ala Ile Ala Thr His Ala Lys Ile  
 346 130 135 140  
 348 Arg Asp  
 349 145  
 351 <210> SEQ ID NO: 21  
 352 <211> LENGTH: 261  
 353 <212> TYPE: PRT  
 354 <213> ORGANISM: Dermatophagoides pteronyssinus  
 356 <400> SEQUENCE: 21  
 358 Met Ile Ile Tyr Asn Ile Leu Ile Val Leu Leu Leu Ala Ile Asn Thr  
 359 1 5 10 15

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 09/03/2004  
PATENT APPLICATION: US/10/809,689 TIME: 11:39:26

Input Set : A:\5538-1010seqlisting.txt  
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**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:22; Xaa Pos. 3,10,16  
Seq#:24; Xaa Pos. 4  
Seq#:43; Xaa Pos. 103  
Seq#:91; Xaa Pos. 5,17  
Seq#:92; Xaa Pos. 39  
Seq#:112; Xaa Pos. 3,28

VERIFICATION SUMMARY  
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Input Set : A:\5538-1010seqlisting.txt  
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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:421 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:22  
L:421 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0  
L:473 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:24  
L:473 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0  
L:1280 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:43  
L:1280 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:96  
L:3610 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:91  
L:3610 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:91 after pos.:0  
M:341 Repeated in SeqNo=91  
L:3634 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:92  
L:3634 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:92 after pos.:32  
L:4955 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:112  
L:4955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112 after pos.:0  
M:341 Repeated in SeqNo=112

**STATISTICS SUMMARY**  
PATENT APPLICATION: US/10/809,689

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TIME: 11:39:26

Input Set : A:\5538-1010seqlisting.txt  
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Application Serial Number: US/10/809,689

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 03-25-2004

Art Unit: IFWO

Software Application: PatentIN3.0

Total Number of Sequences: 124

Total Nucleotides: 0

Total Amino Acids: 24874

Number of Errors: 0

Number of Warnings: 14

Number of Corrections: 2

#### **MESSAGE SUMMARY**

258 W: 6 (Mandatory Feature missing)  
270 C: 1 (Current Application Number differs)  
271 C: 1 (Current Filing Date differs)  
341 W: 8 ((46) "n" or "Xaa" used)